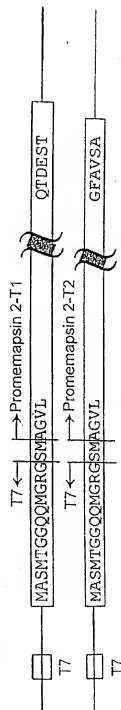


FIG. 1

14 MASMTGGQOM GRGSMAGVLP AHGTQHGI RLPLRSLGGAP LGLRLPRETD  
 36 EEPEEPGRRG SFVEMVDNLR GKSGQGYVE MTVGSPQTL NILVDTGSSN  
 38 FAVGAAPHPF LHRYYQRQLS STYRDLRKGV YVPYTGKWE GELGTDLVSI  
 88 PHGPNVTVRA NIAAITESDK FFINGSNWEG ILGLAYAEIA RPDDSLPEPF  
 138 DSLVKQTHVP NLFSLQLCGA GFPLNQSEVL ASVGSMIIG GIDHSLYTGS  
 188 LWYTPIRREW YVEVIIVRVE INGQDLKMDK KEYNYDKSIV DSGTTNLRLP  
 238 KKVFEAAVKS IKAASSTEF PDGFWLGEQL VCWQAGTTPW NIFPVISLYL  
 288 MGEVTNQSR ITILPQQYLR PVEDVATSD DCYKFAISQS STGTWNGAVI  
 338 MEGFYVVVDR ARKRIGFAVS ACHVHDEFRT AAVEGPFVTL DMEDCGYNIP  
 388 QTDESTILMTI AYVMAAICAL FMLPLCLMVC QWRCLRLCLRQ QHDDFADDIS  
 438 LLK\*



T7 Protein (SEQ ID NO: 3)  
 Memapsin 2T1 and 2T2 Construct (SEQ ID NOS: 32-34)